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RAW SEQUENCE LISTING

DATE: 07/02/2002

PATENT APPLICATION: US/10/045,594B

TIME: 10:16:38

Input Set : A:\DC-0134\Dc-0134.app

Output Set: N:\CRF3\07022002\J045594B.raw

ENTERED

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3 <110> APPLICANT: Taylor, Ronald K.
4   LaPointe, Christian F.
5   Trustees of Dartmouth College
7 <120> TITLE OF INVENTION: COMPOUNDS AND METHODS OF IDENTIFYING COMPOUNDS WHICH
8   INHIBIT A NEW CLASS OF ASPARTYL PROTEASES
10 <130> FILE REFERENCE: DC-0134
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/045,594B
C--> 13 <141> CURRENT FILING DATE: 2002-06-17
15 <150> PRIOR APPLICATION NUMBER: 60/143,355
16 <151> PRIOR FILING DATE: 1999-07-12
18 <160> NUMBER OF SEQ ID NOS: 4
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23 <211> LENGTH: 705
24 <212> TYPE: DNA
25 <213> ORGANISM: Staphylococcus aureus
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30 tgcgcactca aatgggtatga attaatgccg attattagtt ttttattatt aaaagggcga 180
31 tgtcgaaact gtcgaaagcg tatttcoccta acacatttct taggggaaac ctttgcttta 240
32 atacctatcg totttattaa gtatgatttc acatacgtaa atgctacgct atttataact 300
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35 tttatcatta ttagtatgac cacgcatata ttctactttt tatttcgggc atatattggt 480
36 tatggtgacg ttttactaat atctgcactt tctttgtttt tccctctcca attcactatt 540
37 tatgtcattt tatttacatt tgtcattgct ggtttagttg ctttaattac catgatattt 600
38 aagccgatta aactattacc ccttggtcca tttatattta tttccttttt tatcaattca 660
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43 <211> LENGTH: 235
44 <212> TYPE: PRT
45 <213> ORGANISM: Staphylococcus aureus
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51 Tyr Gln Phe Ile Ser Ile Glu Glu Thr Ser Phe Asp Tyr Leu His Arg
52           20           25           30
54 Arg Ser Lys Cys Asp Tyr Cys Asn Ser Ser Leu Lys Trp Tyr Glu Leu
55           35           40           45
57 Met Pro Ile Ile Ser Phe Leu Leu Leu Lys Gly Arg Cys Arg Asn Cys
58           50           55           60
60 Arg Lys Arg Ile Ser Leu Thr His Phe Leu Gly Glu Thr Phe Ala Leu

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61 65          70          75          80
63 Ile Pro Ile Val Phe Ile Lys Tyr Asp Phe Thr Tyr Val Asn Ala Thr
64          85          90          95
66 Leu Phe Ile Thr Thr Tyr Val Phe Leu Leu Ile Phe Thr Met Thr Asp
67          100          105          110
69 Ile Thr Ser Leu Met Leu Asp Cys Arg Leu Ile Ile Ile Tyr Cys Ile
70          115          120          125
72 Val Ser Leu Ser Leu Ser Met Ile Tyr Pro Val Ala Phe Ile Ile Ile
73          130          135          140
75 Ser Met Thr Thr His Ile Phe Tyr Phe Leu Phe Arg Ala Tyr Ile Gly
76 145          150          155          160
78 Tyr Gly Asp Val Leu Leu Ile Ser Ala Leu Ser Leu Phe Phe Pro Leu
79          165          170          175
81 Gln Phe Thr Ile Tyr Val Ile Leu Phe Thr Phe Val Ile Ala Gly Leu
82          180          185          190
84 Val Ala Leu Ile Thr Met Ile Phe Lys Pro Ile Lys Leu Leu Pro Leu
85          195          200          205
87 Val Pro Phe Ile Phe Ile Ser Phe Phe Ile Asn Ser Leu Phe Tyr Asn
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96 <212> TYPE: DNA
97 <213> ORGANISM: Vibrio cholerae
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102 gttgcagaaa aataatggaa tacgtttact tgatcctatt ttcgattgta tcaactgattt 180
103 tgggtagttt tagtaatggt gtgattttatc gtctaccaag aaaaatttta ctaaagaatc 240
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113 catttattca ttggtttttt gctagagtct catgtcggta ttgcttatat atccctcttg 840
114 gaccaagcat tattatctct tttgtaattg tttttttttc aatccgttta atgtgagcta 900
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126      1              5              10              15
128 Gly Ser Phe Ser Asn Val Val Ile Tyr Arg Leu Pro Arg Lys Ile Leu
129              20              25              30
131 Leu Lys Asn His Phe Phe Tyr Asp Ile Asp Ser Asn Arg Ser Met Cys
132              35              40              45
134 Pro Lys Cys Gly Asn Lys Ile Ser Trp Tyr Asp Asn Val Pro Leu Leu
135              50              55              60
137 Ser Tyr Leu Leu Leu His Gly Lys Cys Arg His Cys Asp Glu Lys Ile
138      65              70              75              80
140 Ser Leu Ser Tyr Phe Ile Val Glu Leu Ser Phe Phe Ile Ile Ala Phe
141              85              90              95
143 Pro Ile Tyr Trp Leu Ser Thr Asp Trp Val Asp Ser Phe Val Leu Leu
144              100             105             110
146 Gly Leu Tyr Phe Ile Leu Phe Asn Leu Phe Val Ile Asp Phe Lys Ser
147              115             120             125
149 Met Leu Leu Pro Asn Leu Leu Thr Tyr Pro Ile Phe Met Leu Ala Phe
150              130             135             140
152 Ile Tyr Val Gln Pro Asn Gln Ala Leu Thr Val Glu Ser Ser Ile Ile
153 145              150             155             160
155 Gly Gly Phe Ala Ala Phe Ile Ile Thr Tyr Val Ser Asn Phe Ile Val
156              165             170             175
158 Arg Leu Phe Lys Arg Ile Asp Val Met Gly Gly Gly Asp Ile Lys Leu
159              180             185             190
161 Tyr Thr Ala Ile Gly Thr Leu Ile Gly Val Glu Phe Val Pro Tyr Leu
162              195             200             205
164 Phe Leu Leu Ser Ser Ile Ile Ala Phe Ile His Trp Phe Phe Ala Arg
165              210             215             220
167 Val Ser Cys Arg Tyr Cys Leu Tyr Ile Pro Leu Gly Pro Ser Ile Ile
168 225              230             235             240
170 Ile Ser Phe Val Ile Val Phe Phe Ser Ile Arg Leu Met
171              245             250

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• VERIFICATION SUMMARY

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L:12 M:270 C: Current Application Number differs, Replaced Application Number

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date